Table S3. Oligonucleotides used for the molecular identification and/or characterization of the enteric protist species investigated in the present study.

Target organism	Locus	Oligonucleotide	Sequence (5´–3´)	Reference
Cryptosporidium spp.	*ssu* rRNA	CR-P1	CAGGGAGGTAGTGACAAGAA	(30)
		CR-P2	TCAGCCTTGCGACCATACTC	
		CR-P3	ATTGGAGGGCAAGTCTGGTG	
		CPB-DIAGR	TAAGGTGCTGAAGGAGTAAGG	
Entamoeba histolytica	*ssu* rRNA	Probe	FAM–TCATTGAATGAATTGGCCATTT–MGB	(32)
Entamoeba dispar		Probe	VIC–TTACTTACATAAATTGGCCACTTTG–MGB	
Entamoeba		Ehd-239F	ATTGTCGTGGCATCCTAACTCA	(31)
histolytica/*dispar*		Ehd-88R	GCGGACGGCTCATTATAACA	
Giardia duodenalis	*ssu* rRNA	Probe	FAM–CCCGCGGCGGTCCCTGCTAG–BHQ1	(33)
		Gd-80F	GACGGCTCAGGACAACGGTT	
		Gd-127R	TTGCCAGCGGTGTCCG	
	gdh	GDHeF	TCAACGTYAAYCGYGGYTTCCGT	(34)
		GDHiF	CAGTACACCTCYGCTCTCGG	
		GDHiR	GTTRTCCTTGCACATCTCC	
	bg	G7_F	AAGCCCGACGACCTCACCCGCAGTGC	(35)
		G759_R	GAGGCCGCCCTGGATCTTCGAGACGAC	

		G99_F	GAACGAACGAGATCGAGGTCCG	
		G609_R	CTCGACGAGCTTCGTGTT	
	tpi	AL3543	AAATIATGCCTGCTCGTCG	(36)
		AL3546	CAAACCTTITCCGCAAACC	
		AL3544	CCCTTCATCGGIGGTAACTT	
		AL3545	GTGGCCACCACICCCGTGCC	
Blastocystis spp.	*ssu* rRNA	BhRDr	GAGCTTTTTAACTGCAACAACG	(37)
		RD5	ATCTGGTTGATCCTGCCAGT	
Enterocytozoon bieneusi	ITS	EBITS3	GGTCATAGGGATGAAGAG	(38)
		EBITS4	TTCGAGTTCTTTCGCGCTC	
		EBITS1	GCTCTGAATATCTATGGCT	
		EBITS2.4	ATCGCCGACGGATCCAAGTG	
Balantioides coli	ITS	B5D	GCTCCTACCGATACCGGGT	(39)
		B5RC	GCGGGTCATCTTACTTGATTTC	
Troglodytella spp.	ITS	SSU_end)	AAGGTWTCCGTAGGTGAACCTG	(40)
		LSU_start	TAKTRAYATGCTTAAGTYCAGCG	

bg: β-giardin (bg); *gdh*: Glutamate dehydrogenase; ITS: Internal transcribed spacer; *ssu* rRNA: Small subunit ribosomal RNA; *tpi*: Triose phosphate isomerase.